

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/086,177B
Source: IFW16
Date Processed by STIC: 7/31/06

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IFW16

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/10/086,177B

DATE: 07/31/2006
 TIME: 11:09:32

Input Set : A:\80421-1.APP
 Output Set: N:\CRF4\07312006\J086177B.raw

3 <110> APPLICANT: Tudan, Christopher R.
 4 Merzouk, Ahmed
 5 Arab, Lakhdar
 6 Saxena, Geeta
 7 Eaves, Connie J.
 8 Cashman, Johanne
 9 Clark-Lewis, Ian
 10 Salari, Hassan
 11 University of British Columbia
 12 Chemokine Therapeutics Corporation
 14 <120> TITLE OF INVENTION: CXC Chemokine Receptor 4 Agonist Peptides
 16 <130> FILE REFERENCE: 080421-000100US
 18 <140> CURRENT APPLICATION NUMBER: US 10/086,177B
 19 <141> CURRENT FILING DATE: 2002-02-26
 21 <150> PRIOR APPLICATION NUMBER: CA 2,305,036
 22 <151> PRIOR FILING DATE: 2000-04-12
 24 <150> PRIOR APPLICATION NUMBER: US 60/232,425
 25 <151> PRIOR FILING DATE: 2000-09-14
 27 <150> PRIOR APPLICATION NUMBER: CA 2,335,109
 28 <151> PRIOR FILING DATE: 2001-02-23
 30 <150> PRIOR APPLICATION NUMBER: US 09/835,107
 31 <151> PRIOR FILING DATE: 2001-04-12
 33 <160> NUMBER OF SEQ ID NOS: 214
 35 <170> SOFTWARE: PatentIn version 3.3
 37 <210> SEQ ID NO: 1
 38 <211> LENGTH: 67
 39 <212> TYPE: PRT
 40 <213> ORGANISM: Homo sapiens
 42 <220> FEATURE:
 43 <223> OTHER INFORMATION: human SDF-1alpha
 45 <400> SEQUENCE: 1
 46 Lys Pro Val Ser Leu Ser Tyr Arg Cys Pro Cys Arg Phe Phe Glu Ser
 47 1 5 10 15
 48 His Val Ala Arg Ala Asn Val Lys His Leu Lys Ile Leu Asn Thr Pro
 49 20 25 30
 50 Asn Cys Ala Leu Gln Ile Val Ala Arg Leu Lys Asn Asn Asn Arg Gln
 51 35 40 45
 52 Val Cys Ile Asp Pro Lys Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys
 53 50 55 60
 54 Ala Leu Asn
 55 65
 57 <210> SEQ ID NO: 2
 58 <211> LENGTH: 93

JY 6-7

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59 <212> TYPE: PRT
60 <213> ORGANISM: Homo sapiens
62 <220> FEATURE:
63 <223> OTHER INFORMATION: human SDF-1 precursor, PBSF
65 <400> SEQUENCE: 2
66 Met Asn Ala Lys Val Val Val Val Leu Val Leu Val Leu Thr Ala Leu
67 1 5 10 15
68 Cys Leu Ser Asp Gly Lys Pro Val Ser Leu Ser Tyr Arg Cys Pro Cys
69 20 25 30
70 Arg Phe Phe Glu Ser His Val Ala Arg Ala Asn Val Lys His Leu Lys
71 35 40 45
72 Ile Leu Asn Thr Pro Asn Cys Ala Leu Gln Ile Val Ala Arg Leu Lys
73 50 55 60
74 Asn Asn Asn Arg Gln Val Cys Ile Asp Pro Lys Leu Lys Trp Ile Gln
75 65 70 75 80
76 Glu Tyr Leu Glu Lys Ala Leu Asn Lys Arg Phe Lys Met
77 85 90
79 <210> SEQ ID NO: 3
80 <211> LENGTH: 93
81 <212> TYPE: PRT
82 <213> ORGANISM: Homo sapiens
84 <220> FEATURE:
85 <223> OTHER INFORMATION: human SDF-1beta
87 <400> SEQUENCE: 3
88 Met Asn Ala Lys Val Val Val Val Leu Val Leu Val Leu Thr Ala Leu
89 1 5 10 15
90 Cys Leu Ser Asp Gly Lys Pro Val Ser Leu Ser Tyr Arg Cys Pro Cys
91 20 25 30
92 Arg Phe Phe Glu Ser His Val Ala Arg Ala Asn Val Lys His Leu Lys
93 35 40 45
94 Ile Leu Asn Thr Pro Asn Cys Ala Leu Gln Ile Val Ala Arg Leu Lys
95 50 55 60
96 Asn Asn Asn Arg Gln Val Cys Ile Asp Pro Lys Leu Lys Trp Ile Gln
97 65 70 75 80
98 Glu Tyr Leu Glu Lys Ala Leu Asn Lys Arg Phe Lys Met
99 85 90
101 <210> SEQ ID NO: 4
102 <211> LENGTH: 17
103 <212> TYPE: PRT
104 <213> ORGANISM: Artificial Sequence
106 <220> FEATURE:
107 <223> OTHER INFORMATION: synthetic CXCR4 agonist SDF-1(1-17), CTCE9902
109 <400> SEQUENCE: 4
110 Lys Pro Val Ser Leu Ser Tyr Arg Cys Pro Cys Arg Phe Phe Glu Ser
111 1 5 10 15
112 His
115 <210> SEQ ID NO: 5
116 <211> LENGTH: 6
117 <212> TYPE: PRT

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118 <213> ORGANISM: Artificial Sequence
 120 <220> FEATURE:
 121 <223> OTHER INFORMATION: CXCR4 agonist sequence motif within 20 amino acids
 122 of the N-terminus
 124 <400> SEQUENCE: 5
 125 Arg Phe Phe Glu Ser His
 126 1 5
 128 <210> SEQ ID NO: 6
 129 <211> LENGTH: 9
 130 <212> TYPE: PRT
 131 <213> ORGANISM: Artificial Sequence
 133 <220> FEATURE:
 134 <223> OTHER INFORMATION: synthetic SDF-1 peptide analogue CXCR4 agonist
 136 <400> SEQUENCE: 6
 137 Lys Pro Val Ser Leu Ser Tyr Arg Cys
 138 1 5
 140 <210> SEQ ID NO: 7
 141 <211> LENGTH: 9
 142 <212> TYPE: PRT
 143 <213> ORGANISM: Artificial Sequence
 145 <220> FEATURE:
 146 <223> OTHER INFORMATION: synthetic CXCR4 agonist SDF-1(1-9)-2-C9/C9-cysteine dimer,
 147 CTCE9901
 149 <220> FEATURE:
 150 <221> NAME/KEY: MISC_FEATURE
 151 <222> LOCATION: (7)...(7)
 152 <223> OTHER INFORMATION: dimerised by formation of a disulfide bond between two Cys
 153 residues in position 7 of two SEQ ID NO:7 peptides
 155 <400> SEQUENCE: 7
 156 Lys Pro Val Ser Leu Ser Tyr Arg Cys
 157 1 5
 159 <210> SEQ ID NO: 8
 160 <211> LENGTH: 10
 161 <212> TYPE: PRT
 162 <213> ORGANISM: Artificial Sequence
 164 <220> FEATURE:
 165 <223> OTHER INFORMATION: portion of synthetic CXCR4 agonist SDF-1(1-9)-2
 166 (Compound #3)
 168 <220> FEATURE:
 169 <221> NAME/KEY: MOD_RES
 170 <222> LOCATION: (10)...(10)
 171 <223> OTHER INFORMATION: Xaa = Lys whose epsilon amino group forms a covalent amide
 172 bond with the alpha amino group of Cys at position 9 of
 173 KPVSLSYRC (SEQ ID NO:9), thereby forming a dimer
 175 <400> SEQUENCE: 8
 W--> 176 Lys Pro Val Ser Leu Ser Tyr Arg Cys Xaa
 177 1 5 10
 179 <210> SEQ ID NO: 9
 180 <211> LENGTH: 9

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181 <212> TYPE: PRT
182 <213> ORGANISM: Artificial Sequence
184 <220> FEATURE:
185 <223> OTHER INFORMATION: portion of synthetic CXCR4 agonist SDF-1(1-9)-2
186 (Compound #3)
188 <220> FEATURE:
189 <221> NAME/KEY: MOD_RES
190 <222> LOCATION: (9)..(9)
191 <223> OTHER INFORMATION: Xaa = Cys whose alpha amino group forms a covalent amide
192 bond with the epsilon amino group of Lys at position 10
193 of KPVSLSYRCX (SEQ ID NO:8), thereby forming a dimer
195 <400> SEQUENCE: 9
W--> 196 Lys Pro Val Ser Leu Ser Tyr Arg Xaa
197 1 5
199 <210> SEQ ID NO: 10
200 <211> LENGTH: 9
201 <212> TYPE: PRT
202 <213> ORGANISM: Artificial Sequence
204 <220> FEATURE:
205 <223> OTHER INFORMATION: portion of synthetic CXCR4 agonist dimer of SDF-1 amino
206 acids 1-8
208 <220> FEATURE:
209 <221> NAME/KEY: MOD_RES
210 <222> LOCATION: (9)..(9)
211 <223> OTHER INFORMATION: Xaa = Lys whose epsilon amino group forms a covalent amide
212 bond with the alpha amino group of Arg at position 8 of
213 KPVSLSYX (SEQ ID NO:11), thereby forming a dimer
215 <400> SEQUENCE: 10
W--> 216 Lys Pro Val Ser Leu Ser Tyr Arg Xaa
217 1 5
219 <210> SEQ ID NO: 11
220 <211> LENGTH: 8
221 <212> TYPE: PRT
222 <213> ORGANISM: Artificial Sequence
224 <220> FEATURE:
225 <223> OTHER INFORMATION: portion of synthetic CXCR4 agonist dimer of SDF-1 amino
226 acids 1-8
228 <220> FEATURE:
229 <221> NAME/KEY: MOD_RES
230 <222> LOCATION: (8)..(8)
231 <223> OTHER INFORMATION: Xaa = Arg whose alpha amino group forms a covalent amide
232 bond with the epsilon amino group of Lys at position 9
233 of KPVSLSYRX (SEQ ID NO:10), thereby forming a dimer
235 <400> SEQUENCE: 11
W--> 236 Lys Pro Val Ser Leu Ser Tyr Xaa
237 1 5
239 <210> SEQ ID NO: 12
240 <211> LENGTH: 30
241 <212> TYPE: PRT

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242 <213> ORGANISM: Artificial Sequence
244 <220> FEATURE:
245 <223> OTHER INFORMATION: synthetic CXCR4 agonist SDF-1(1-14) - (G) -3-SDF-1(55-67)
246 acid
248 <220> FEATURE:
249 <221> NAME/KEY: MISC_FEATURE
250 <222> LOCATION: (17)..(17)
251 <223> OTHER INFORMATION: Gly in position 17 may be present or absent
253 <400> SEQUENCE: 12
254 Lys Pro Val Ser Leu Ser Tyr Arg Cys Pro Cys Arg Phe Phe Gly Gly
255 1 5 10 15
256 Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
257 20 25 30
259 <210> SEQ ID NO: 13
260 <211> LENGTH: 31
261 <212> TYPE: PRT
262 <213> ORGANISM: Artificial Sequence
264 <220> FEATURE:
265 <223> OTHER INFORMATION: synthetic CXCR4 agonist SDF-1(1-14) - (G) -4-SDF-1(55-67)
266 acid, CTCE0013
268 <220> FEATURE:
269 <221> NAME/KEY: MISC_FEATURE
270 <222> LOCATION: (17)..(18)
271 <223> OTHER INFORMATION: Gly in positions 17 and/or 18 may independently be
272 present or absent
274 <400> SEQUENCE: 13
275 Lys Pro Val Ser Leu Ser Tyr Arg Cys Pro Cys Arg Phe Phe Gly Gly
276 1 5 10 15
277 Gly Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn
278 20 25 30
280 <210> SEQ ID NO: 14
281 <211> LENGTH: 30
282 <212> TYPE: PRT
283 <213> ORGANISM: Artificial Sequence
285 <220> FEATURE:
286 <223> OTHER INFORMATION: synthetic CXCR4 agonist SDF-1(1-14) - (G) -3-SDF-1(55-67)
287 amide
289 <220> FEATURE:
290 <221> NAME/KEY: MISC_FEATURE
291 <222> LOCATION: (17)..(17)
292 <223> OTHER INFORMATION: Gly in position 17 may be present or absent
294 <220> FEATURE:
295 <221> NAME/KEY: MOD_RES
296 <222> LOCATION: (30)..(30)
297 <223> OTHER INFORMATION: AMIDATION
299 <400> SEQUENCE: 14
300 Lys Pro Val Ser Leu Ser Tyr Arg Cys Pro Cys Arg Phe Phe Gly Gly
301 1 5 10 15
302 Gly Leu Lys Trp Ile Gln Glu Tyr Leu Glu Lys Ala Leu Asn

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:8; Xaa Pos. 10
Seq#:9; Xaa Pos. 9
Seq#:10; Xaa Pos. 9
Seq#:11; Xaa Pos. 8
Seq#:31; Xaa Pos. 9
Seq#:32; Xaa Pos. 8
Seq#:35; Xaa Pos. 2,5
Seq#:36; Xaa Pos. 2,5
Seq#:39; Xaa Pos. 20
Seq#:40; Xaa Pos. 20
Seq#:41; Xaa Pos. 28
Seq#:42; Xaa Pos. 28
Seq#:43; Xaa Pos. 20
Seq#:44; Xaa Pos. 20
Seq#:45; Xaa Pos. 28
Seq#:46; Xaa Pos. 28
Seq#:50; Xaa Pos. 2,5
Seq#:51; Xaa Pos. 2,5,14
Seq#:52; Xaa Pos. 1
Seq#:53; Xaa Pos. 2,5
Seq#:54; Xaa Pos. 2,5,14
Seq#:55; Xaa Pos. 1
Seq#:58; Xaa Pos. 14
Seq#:59; Xaa Pos. 1
Seq#:61; Xaa Pos. 14
Seq#:62; Xaa Pos. 1
Seq#:64; Xaa Pos. 14
Seq#:65; Xaa Pos. 1
Seq#:68; Xaa Pos. 1
Seq#:70; Xaa Pos. 14
Seq#:71; Xaa Pos. 1
Seq#:73; Xaa Pos. 14
Seq#:74; Xaa Pos. 1
Seq#:76; Xaa Pos. 14
Seq#:77; Xaa Pos. 1
Seq#:79; Xaa Pos. 14
Seq#:80; Xaa Pos. 1
Seq#:82; Xaa Pos. 14
Seq#:83; Xaa Pos. 1
Seq#:85; Xaa Pos. 14
Seq#:86; Xaa Pos. 1
Seq#:88; Xaa Pos. 14
Seq#:89; Xaa Pos. 1
Seq#:91; Xaa Pos. 14

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Seq#:92; Xaa Pos. 1
Seq#:94; Xaa Pos. 14
Seq#:95; Xaa Pos. 1
Seq#:97; Xaa Pos. 14
Seq#:98; Xaa Pos. 1
Seq#:100; Xaa Pos. 14
Seq#:101; Xaa Pos. 1

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Input Set : A:\80421-1.APP
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L:176 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0
L:196 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
L:216 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
L:236 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0
L:746 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 after pos.:0
L:767 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:0
L:773 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (33) SEQUENCE:
L:778 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (34) SEQUENCE:
L:824 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:0
L:873 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:0
L:966 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39 after pos.:0
L:1003 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40 after pos.:16
L:1035 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41 after pos.:16
L:1072 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:42 after pos.:16
L:1104 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:43 after pos.:16
L:1141 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:44 after pos.:16
L:1173 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45 after pos.:16
L:1210 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:46 after pos.:16
L:1341 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:50 after pos.:0
L:1382 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51 after pos.:0
L:1413 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:52 after pos.:0
L:1460 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:53 after pos.:0
L:1501 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:54 after pos.:0
L:1532 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:55 after pos.:0
L:1605 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:58 after pos.:0
L:1635 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59 after pos.:0
L:1689 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:61 after pos.:0
L:1719 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:62 after pos.:0
L:1763 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:64 after pos.:0
L:1793 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:65 after pos.:0
L:1882 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:68 after pos.:0
L:1926 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:70 after pos.:0
L:1956 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71 after pos.:0
L:2010 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:73 after pos.:0
L:2040 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:74 after pos.:0
L:2084 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:76 after pos.:0
L:2114 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:77 after pos.:0
L:2168 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:79 after pos.:0
L:2198 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:80 after pos.:0
L:2242 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:82 after pos.:0
L:2272 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:83 after pos.:0
L:2326 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:85 after pos.:0
L:2356 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:86 after pos.:0
L:2400 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:88 after pos.:0
L:2430 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:89 after pos.:0
L:2484 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:91 after pos.:0
L:2514 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:92 after pos.:0
L:2558 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:94 after pos.:0

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L:2588 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:95 after pos.:0
L:2642 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:97 after pos.:0
L:2672 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:98 after pos.:0
L:2716 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:100 after pos.:0